



00670658.TXT

## SEQUENCE LISTING

<110> Chu, Rea-Min  
Lin, Ching-Yi  
Hsiao, Ya-Wen  
Liao, Kuang-Wen

<120> COMPLEX IMMUNO-GENE MEDICAL COMPOSITION FOR INHIBITING TUMOR CELLS

<130> P/741-176

<140> 10/798,096

<141> 2004-03-11

<160> 4

<170> PatentIn version 3.2

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<221> CDS

<222> (1)..(636)

<223> IL-6

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Gly	Leu	Leu	Leu	Val	Leu	Pro	Ala	Ala	Phe	Pro	Ala	Pro	Val	Pro	Pro	
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gga	gaa	gat	tcc	aaa	gat	gta	gcc	gcc	cca	cac	aga	cag	cca	ctc	acc	144
Gly	Glu	Asp	Ser	Lys	Asp	Val	Ala	Ala	Pro	His	Arg	Gln	Pro	Leu	Thr	
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tct	tca	gaa	cga	att	gac	aaa	caa	att	cgg	tac	atc	ctc	gac	ggc	atc	192
Ser	Ser	Glu	Arg	Ile	Asp	Lys	Gln	Ile	Arg	Tyr	Ile	Leu	Asp	Gly	Ile	
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tca	gcc	ctg	aga	aag	gag	aca	tgt	aac	aag	agt	aac	atg	tgt	gaa	agc	240
Ser	Ala	Leu	Arg	Lys	Glu	Thr	Cys	Asn	Lys	Ser	Asn	Met	Cys	Glu	Ser	
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agc	aaa	gag	gca	ctg	gca	gaa	aac	aac	ctg	aac	ctt	cca	aag	atg	gct	288
Ser	Lys	Glu	Ala	Leu	Ala	Glu	Asn	Asn	Leu	Asn	Leu	Pro	Lys	Met	Ala	
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gaa	aaa	gat	gga	tgc	ttc	caa	tct	gga	ttc	aat	gag	gag	act	tgc	ctg	336
Glu	Lys	Asp	Gly	Cys	Phe	Gln	Ser	Gly	Phe	Asn	Glu	Glu	Thr	Cys	Leu	
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gtg	aaa	atc	atc	act	ggt	ctt	ttg	gag	ttt	gag	gta	tac	cta	cag	tac	384
Val	Lys	Ile	Ile	Thr	Gly	Leu	Leu	Glu	Phe	Glu	Val	Tyr	Leu	Gln	Tyr	
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ctc	cag	aac	aga	ttt	gag	agt	agt	gag	gaa	caa	gcc	aga	gct	gtg	cag	432

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Met	Ser	Thr	Lys	Val	Leu	Ile	Gln	Phe	Leu	Gln	Lys	Lys	Ala	Lys	Asn	160	
145				150					155								
cta	gat	gca	ata	acc	cct	gac	cca	acc	aca	aat	gcc	agc	ctg	ctg	528		
Leu	Asp	Ala	Ile	Thr	Pro	Asp	Pro	Thr	Thr	Asn	Ala	Ser	Leu	Leu	175		
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acg	aag	ctg	cag	gca	cag	aac	cag	tgg	ctg	cag	gac	atg	aca	act	cat	576	
Thr	Lys	Leu	Gln	Ala	Gln	Asn	Gln	Trp	Leu	Gln	Asp	Met	Thr	Thr	His	190	
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ctc	att	ctg	cgc	agc	ttt	aag	gag	ttc	ctg	cag	tcc	agc	ctg	agg	gct	624	
Leu	Ile	Leu	Arg	Ser	Phe	Lys	Glu	Phe	Leu	Gln	Ser	Ser	Leu	Arg	Ala	200	
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1				5				10									
caa	tct	atg	cat	att	gat	gct	act	tta	tat	acg	gaa	agt	gat	gtt	cac	96	
Gln	Ser	Met	His	Ile	Asp	Ala	Thr	Leu	Tyr	Thr	Glu	Ser	Asp	Val	His	20	
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ccc	agt	tgc	aaa	gta	aca	gca	atg	aag	tgc	ttt	ctc	ttg	gag	tta	caa	144	
Pro	Ser	Cys	Lys	Val	Thr	Ala	Met	Lys	Cys	Phe	Leu	Leu	Glu	Leu	Gln		

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aca gaa tct gga tgc aaa gaa tgt gag gaa ctg gag gaa aaa aat att Thr Glu Ser Gly Cys Lys Glu Cys Glu Glu Leu Glu Glu Lys Asn Ile 85 90 95			288
aaa gaa ttt ttg cag agt ttt gta cat att gtc caa atg ttc atc aac Lys Glu Phe Leu Gln Ser Phe Val His Ile Val Gln Met Phe Ile Asn 100 105 110			336
act tct Thr Ser			342
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gtc aca aac agt aac tgg gtg aat gta ata agt gat ttg aaa aaa att Val Thr Asn Ser Asn Trp Val Asn Val Ile Ser Asp Leu Lys Lys Ile 20 25 30			96
gaa gat ctt att caa tct atg cat att gat gct act tta tat acg gaa Glu Asp Leu Ile Gln Ser Met His Ile Asp Ala Thr Leu Tyr Thr Glu 35 40 45			144
agt gat gtt cac ccc agt tgc aaa gta aca gca atg aag tgc ttt ctc Ser Asp Val His Pro Ser Cys Lys Val Thr Ala Met Lys Cys Phe Leu 50 55 60			192
ttg gag tta caa gtt att tca ctt gag tcc gga gat gca agt att cat Leu Glu Leu Gln Val Ile Ser Leu Glu Ser Gly Asp Ala Ser Ile His 65 70 75 80			240
gat aca gta gaa aat ctg atc atc cta gca aac aac agt ttg tct tct Asp Thr Val Glu Asn Leu Ile Ile Leu Ala Asn Asn Ser Leu Ser Ser 85 90 95			288
aat ggg aat gta aca gaa tct gga tgc aaa gaa tgt gag gaa ctg gag Asn Gly Asn Val Thr Glu Ser Gly Cys Lys Glu Cys Glu Glu Leu Glu 100 105 110			336
gaa aaa aat att aaa gaa ttt ttg cag agt ttt gta cat att gtc caa Glu Lys Asn Ile Lys Glu Phe Leu Gln Ser Phe Val His Ile Val Gln 115 120 125			384

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atg ttc atc aac act tct  
Met Phe Ile Asn Thr Ser  
130

402